Amendments to the Specification:

Please replace the paragraph beginning at page 23, line 20, with the following:

--To verify the microarray data, six genes were selected and their expression levels examined in 13 additional samples (7 adenomas and 6 carcinomas) by means of real-time quantitative RT-PCR (TaqMan PCR, Perkin-Elmer), using a 7700 Sequence Detector (Perkin-Elmer). Each single-stranded cDNA was reverse-transcribed from amplified RNA and diluted for subsequent PCR amplification. Malate dehydrogenase 1 (*MDH1*) served as a relative quantitative control since it showed the smallest Cy3/Cy5 fluctuations in 100 hybridizations. Each PCR was carried out in a 25-μl volume and amplified for 10 min at 95^{O}C for activation of AmpliTaq Gold™, followed by 40 cycles of 95^{O}C for 15 s and 60^{O}C for 1 min. The genes and sequences of the primers and probes used for quantitative RT-PCR are listed in Table A below (SEQ ID NOS:1-21).--

Please cancel the present "SEQUENCE LISTING", pages 1/13-13/13, and insert therefor the accompanying paper copy of the Substitute Sequence Listing, page numbers 1 to 6, at the end of the application.